

AMENDED CLAIMS

- 1. (Amended) A polynucleotide comprising a non-naturally occurring HCV sequence that is capable of productive replication in a host cell, or is capable of being transcribed into a non-naturally occurring HCV sequence that is capable of productive replication in a host cell, wherein the HCV sequence comprises, from 5' to 3' on the positive-sense nucleic acid, a functional 5' non-translated region (5' NTR); one or more protein coding regions, including at least one polyprotein coding region that is capable of replicating HCV RNA; and a functional HCV 3' non-translated region (3' NTR) and wherein said polypeptide further comprises an adaptive mutation.
- 3. (Amended) The polynucleotide of claim 1, having a transfection efficiency into mammalian cells of greater than 0.01%.
- 7. (Amended) The polynucleotide of claim 1, wherein the polynucleotide is capable of replication in a non-hepatic cell.
- 9. (Amended) The polynucleotide of claim 1, wherein the HCV is impaired in its ability to cause disease, establish chronic infections, trigger autoimmune responses, and transform cells.
- 10. (Amended) The polynucleotide of claim 1, wherein the polyprotein region comprises an NS5A gene that is not a wild-type NS5A gene.
- 29. (Amended) The polynucleotide of claim 1, wherein the transfection efficiency into mammalian cells is about 6%
- 65. (Amended) The vector of claim 64, wherein the adaptive mutation comprises a mutation in the NS5A gene.